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<110> Reactive Surfaces
McDaniel, C. Steven

<120> Recombinant Organophosphorus Acid Anhydride and Methods of Use

<130> RACT-00200

<140> Unknown
<141> 2002-12-28

<150> 07/928,540
<151> 1992-08-13

<150> 08/252,384
<151> 1994-06-01

<150> 07/344,258
<151> 1989-04-27

<160> 1

<170> PatentIn version 3.2

<210> 1
<211> 337
<212> PRT
<213> Pseudomonas aeruginosa

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Ser Ser Ala Gly Phe Leu Arg Ala Trp Pro Glu Phe Phe Gly Ser Arg
35 40 45

Lys Ala Leu Ala Glu Lys Ala Val Arg Gly Leu Arg Arg Ala Arg Ala
50 55 60

BB
Conf

Ala Gly Val Arg Thr Ile Val Asp Val Ser Thr Phe Asp Ile Gly Arg
65 70 75 80

Asp Val Ser Leu Leu Ala Glu Val Ser Arg Ala Ala Asp Val His Ile
85 90 95

Val Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu
100 105 110

Arg Ser Val Glu Glu Leu Thr Gln Phe Phe Leu Arg Glu Ile Gln Tyr
115 120 125

Gly Ile Glu Asp Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr
130 135 140

Thr Gly Lys Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala
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Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala
165 170 175

Ser Gln Arg Asp Gly Glu Gln Gln Ala Ala Ile Phe Glu Ser Glu Gly
180 185 190

Leu Ser Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp Asp
195 200 205

Leu Ser Tyr Leu Thr Ala Leu Ala Ala Arg Gly Tyr Leu Ile Gly Leu
210 215 220

Asp His Ile Pro His Ser Ala Ile Gly Leu Gln Asp Asn Ala Ser Ala
225 230 235 240

Ser Ala Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile
245 250 255

Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln Ile Leu Val Ser Asn
260 265 270

Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val
275 280 285

B3
Conclude
Met Asp Arg Val Asn Pro Asp Gly Met Ala Phe Ile Pro Leu Arg Val
290 295 300

Ile Pro Phe Leu Arg Glu Lys Gly Val Pro Gln Glu Thr Leu Ala Gly
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Ile Thr Val Thr Asn Pro Ala Arg Phe Leu Ser Pro Thr Leu Arg Ala
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Ser

SEQUENCE LISTING

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<110> Reactive Surfaces, Ltd.

<120> Recombinant Organophosphorus Acid Anhydride and Methods of Use

<130> RACT-00200

<140> Unknown

<141> 2003-01-02

<150> 07/928,540

<151> 1992-08-13

<150> 08/252,384

<151> 1994-06-01

<150> 07/344,258

<151> 1989-04-27

<160> 2

<170> PatentIn version 3.1

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5

10

15

aca atc tct gaa gcg ggt ttc aca ctg act cac gag cac atc tgc ggc
96

Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu His Ile Cys Gly

20

25

30

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144
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35

40

45

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192
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50

55

60

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240
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65

70

75

80

gac gtc agt tta ttg gcc gag gtt tcg cgg gct gcc gac gtt cat atc
288
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85

90

95

gtg gcg qcq acc ggc ttg tgg ttc gac ccc cca ctt tag atg cga ttg
336
Val Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu

100

105

110

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384
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115

120

125

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432
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130

135

140

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Thr Gly Lys Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala

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155

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165

170

175

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185

190

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624

Leu Ser Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp Asp

195

200

205

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672

Leu Ser Tyr Leu Thr Ala Leu Ala Ala Arg Gly Tyr Leu Ile Gly Leu

210

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225

230

235

240

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Ser Ala Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile

245

250

255

aag gcg ctc atc gac caa ggc tac atg aaa caa atc ctc gtt tcg aat
816

Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln Ile Leu Val Ser Asn

260

265

270

gac tgg ctg ttc ggg ttt tcg agc tat gtc acc aac atc atg gac gtg
864

Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val

275

280

285

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912

Met Asp Arg Val Asn Pro Asp Gly Met Ala Phe Ile Pro Leu Arg Val

290

295

300

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Ile Pro Phe Val Arg Glu Lys Gly Val Pro Gln Glu Thr Leu Ala Gly

305

310

315

320

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Ile Thr Val Thr Asn Pro Ala Arg Phe Tyr Val Thr Asp Leu Ala Gly

325

330

335

gtc atg

1014

Val

<210> 2
<211> 337
<212> PRT
<213> Pseudomonas diminuta

<400> 2

Met Ser Ile Gly Thr Gly Asp Arg Ile Asn Thr Val Arg Gly Pro Ile
1 5 10 15

Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu His Ile Cys Gly
20 25 30

Ser Ser Ala Gly Phe Leu Arg Ala Trp Pro Glu Phe Phe Gly Ser Arg
35 40 45

Lys Ala Leu Ala Glu Lys Ala Val Arg Gly Leu Arg Arg Ala Arg Ala
50 55 60

Ala Gly Val Arg Thr Ile Val Asp Val Ser Thr Phe Asp Ile Gly Arg
65 70 75 80

Asp Val Ser Leu Leu Ala Glu Val Ser Arg Ala Ala Asp Val His Ile
85 90 95

Val Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu
100 105 110

Arg Ser Val Glu Glu Leu Thr Gln Phe Phe Leu Arg Glu Ile Gln Tyr
115 120 125

Gly Ile Glu Asp Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr
130 135 140

Thr Gly Lys Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala
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Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala

165

170

175

Ser Gln Arg Asp Gly Glu Gln Gln Ala Ala Ile Phe Glu Ser Glu Gly
180 185 190

Leu Ser Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp Asp
195 200 205

*PB
Conclude*
Leu Ser Tyr Leu Thr Ala Leu Ala Ala Arg Gly Tyr Leu Ile Gly Leu
210 215 220

Asp His Ile Pro His Ser Ala Ile Gly Leu Glu Asp Asn Ala Ser Ala
225 230 235 240

Ser Ala Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile
245 250 255

Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln Ile Leu Val Ser Asn
260 265 270

Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val
275 280 285

Met Asp Arg Val Asn Pro Asp Gly Met Ala Phe Ile Pro Leu Arg Val
290 295 300

Ile Pro Phe Val Arg Glu Lys Gly Val Pro Gln Glu Thr Leu Ala Gly
305 310 315 320

Ile Thr Val Thr Asn Pro Ala Arg Phe Tyr Val Thr Asp Leu Ala Gly
325 330 335

val

Protein Comparisons Percent Identity

DNA Comparisons Percent Identity

McDaniel vs Wild Lab	57.8%
McDaniel vs Serdar	59.5%
Serdar vs Wild Lab	100%

McDaniel vs Wild Lab	88.0%
McDaniel vs Serdar	83.7%
Serdar vs Wild Lab	99.6%

Decoration 'Decoration #1': Shaded (with black at 10% fill) residues that differ from McDaniel.pro.

Decoration 'Decoration #2': Box residues that differ from McDaniel.pro.

1 ATGCAAACGAGAAGGGTTGTGCTC/ [CTGCAGCCGC---GAGAACTCT] McDaniel P diminu
 2 ATGCAAACGAGAAGGGTTGTGCTCAAGTGTGCCGCCGC [CTGCAGAACTCT] Wild lab opd.seq
 1 ATGCAAACGAGAAGGGTTGTGCTCAAGTGTGCCGCCGC [CTGCAGAACTCT] Sordar.seq
 48 GCTCGCGGGCCCTGGCTGGGTGCCGA-CGTCGCTGGATCCCATCGGCACAG McDaniel P diminu
 4 [CTGCAGCTGGATCCGCACAG] TCGATCCGCACAG Wild lab opd.seq
 51 GCTCGCGCCGGCTGGCTGGATCGCGAG[CTGCAGCTGGATCGCGACAG] Sordar.seq
 97 CGGATCCGATCAATAAC-GTGCGCG-TGCTATCACAAATCTCTGAAGGGT McDaniel P diminu
 17 GGGAT[CTGATCAATAAC] GTGCGCG-TGCTATCACAAATCTCTGAAGGGT Wild lab opd.seq
 101 GCGAT[CTGATCAATAAC] GTGCGCG-TGCTATCACAAATCTCTGAAGGGT Sordar.seq
 145 TTGACACTGACTGACGAGAACATCT-CGGCAGGCTCGGACGGATTCTTGC G McDaniel P diminu
 67 TTGACACTGACTCACCAACATCT[CTGGCAGCTCGGCAGGATTCTTGC] Wild lab opd.seq
 151 TTGACACTGACTCACGAGAACATCT[CTGGCAGCTCGGCAGGATTCTTGC] Sordar.seq
 194 TGCTTGGCCAGAGTTCTTGGCTAGC-CGCAAAAGCTCTAGCGGCGAAAAGCTG McDaniel P diminu
 117 TGCTTGGCCAGAGTTCTTGGCTAGC[CTGGCGAAAAGCTG] Wild lab opd.seq
 201 TGCTTGGCCAGAGTTCTTGGCTAGC[CTGGCGAAAAGCTG] Sordar.seq
 243 TGACAGGATTGGCC---GGCAGAGCGCTGGCGAACATTGTCGAT McDaniel P diminu
 167 TGAGACAGGATTGGCC[CTGGCGAGAGCGCTGGCGAACATTGTCGAT] Wild lab opd.seq
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 301 GTCTCGACTTTGATATCGCTGGCGAACCTGCAAGTTATTGGCGGAGGTTC Sordar.seq
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 267 GGGGCCAGGGCGACTATATC[CTGGCGGCCAGCGCCCTTGTGGTGGCGACC] Wild lab opd.seq
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 784 GCGAGTGCGATCACCGCCCTCGGGCATCCGGTGGCAACACGGGCGCTC McDaniel P diminu
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 846 CCTGATCAAGGGCGTCATCGACCAAGGGTACATGAAACAAATCTCTGTT Sordar.seq
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 996 GATCCCATTCTACGAGAGAAGGGCGTCACAGGAAACGGCTGGCAAGGCA Sordar.seq
 1032 TCACTGTCAGTAACCCGGCGCGCGATCTGTGTCACCGACTTCTCCCTGCG--- McDaniel P diminu
 962 TCACTGTCAGTAACCCGGCGCGCGATCTGTGTCACCGACTTCTCCCTGCG--- Wild lab opd.seq
 1046 TCACTGTCAGTAACCCGGCGCGCGATCTGTGTCACCGACTTCTCCCTGCG--- Sordar.seq
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 1085 ATGA
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 Wild lab opd.seq
 Sordar.seq

Decoration 'Decoration #1': Shaded (with black as 10% fill) residues that differ from McDaniel P diminuta.seq.

Decoration 'Decoration #2': Box residues that differ from McDaniel P diminuta.seq.

MAY-21-2003 WED 10:15 AM

MCDANIEL AND ASSOCIATES

5124728181

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Works Cited

McDaniel P. diminuta :

McDaniel,C.S., Harper,L.L. and Wild,J.R., Cloning and sequencing of a plasmid-borne gene (opd) encoding a phosphotriesterase, J. Bacteriol. 170 (5), 2306-2311 (1988). Gene Bank Accession Number: M20392

Oph-lab RC: Wild lab DNA sequence

Serdar: Serdar Sequence obtained from United States Patent and Trademark Office (uspto.gov) Patent Number 5,484,728

MAY-21-2003 WED 10:16 AM

MCDANIEL AND ASSOCIATES

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